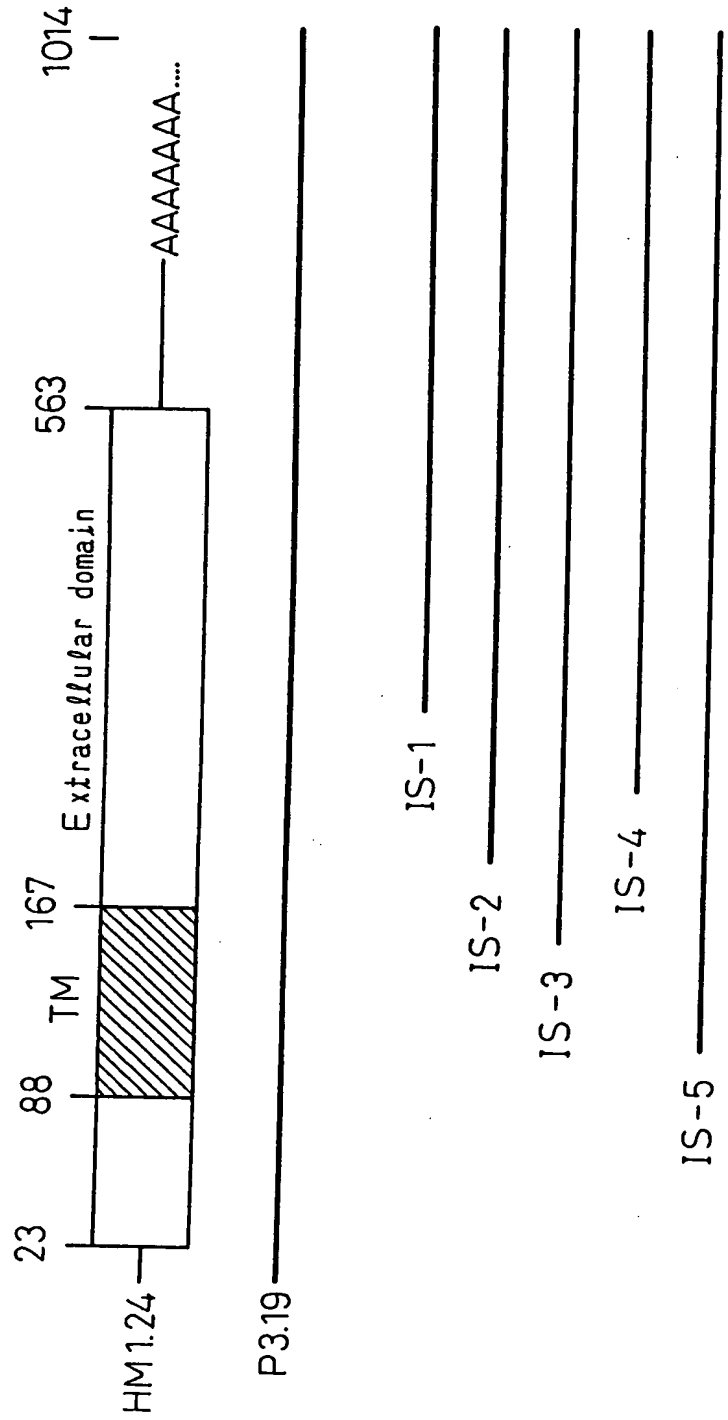
Fig. 14

```
GAATTCGGCACGAGGGATCTGGATGGCATCTACTTCGTATGACTATTGCAGAGTGCCCAT   60
                        M  A  S  T  S  Y  D  Y  C  R  V  P  M   13
GGAAGACGGGGATAAGCGCTGTAAGCTTCTGCTGGGGATAGGAATTCTGGTGCTCCTGAT  120
 E  D  G  D  K  R  C  K  L  L  L  G  I  G  I  L  V  L  L  I   33
CATCGTGATTCTGGGGGTGCCCTTGATTATCTTCACCATCAAGGCCAACAGCGAGGCCTG  180
 I  V  I  L  G  V  P  L  I  I  F  T  I  K  A  N  S  E  A  C   53
CCGGGACGGCCTTCGGGCAGTGATGGAGTGTCGCAATGTCACCCATCTCCTGCAACAAGA  240
 R  D  G  L  R  A  V  M  E  C  R  N  V  T  H  L  L  Q  Q  E   73
GCTGACCGAGGCCCAGAAGGGCTTTCAGGATGTGGAGGCCCAGGCCGCCACCTGCAACCA  300
 L  T  E  A  Q  K  G  F  Q  D  V  E  A  Q  A  A  T  C  N  H   93
CACTGTGATGGCCCTAATGGCTTCCCTGGATGCAGAGAAGGCCCAAGGACAAAAGAAAGT  360
 T  V  M  A  L  M  A  S  L  D  A  E  K  A  Q  G  Q  K  K  V  113
GGAGGAGCTTGAGGGAGAGATCACTACATTAAACCATAAGCTTCAGGACGCGTCTGCAGA  420
 E  E  L  E  G  E  I  T  T  L  N  H  K  L  Q  D  A  S  A  E  133
GGTGGAGCGACTGAGAAGAGAAAACCAGGTCTTAAGCGTGAGAATCGCGGACAAGAAGTA  480
 V  E  R  L  R  R  E  N  Q  V  L  S  V  R  I  A  D  K  K  Y  153
CTACCCCAGCTCCCAGGACTCCAGCTCCGCTGCGGCGCCCCAGCTGCTGATTGTGCTGCT  540
 Y  P  S  S  Q  D  S  S  S  A  A  A  P  Q  L  L  I  V  L  L  173

GGGCCTCAGCGCTCTGCTGCAGTGAGATCCCAGGAAGCTGGCACATCTTGGAAGGTCCGT  600
 G  L  S  A  L  L  Q  *  (SEQ ID NO:26)                       180
CCTGCTCGGCTTTTCGCTTGAACATTCCCTTGATCTCATCAGTTCTGAGCGGGTCATGGG  660
GCAACACGGTTAGCGGGGAGAGCACGGGGTAGCCGGAGAAGGGCCTCTGGAGCAGGTCTG  720
GAGGGGCCATGGGGCAGTCCTGGGTGTGGGGACACAGTCGGGTTGACCCAGGGCTGTCTC  780
CCTCGAGAGCCTCCCTCCGGACAATGAGTCCCCCCTCTTGTCTCCCACCCTGAGATTGGG  840
CATGGGGTGCGGTGTGGGGGGCATGTGCTGCCTGTTGTTATGGGTTTTTTTGCGGGGGG  900
GGTTGCTTTTTTCTGGGGTCTTTGAGCTCCAAAAAAATAAACACTTCCTTTGAGGGAGAG  960
CACACCTTAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAATTCGGGCGGCCGCCA       1014
(SEQ ID NO:16)
```

REPLACEMENT SHEET

Fig. 15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AAAAAAA....
563
Extracellular domain
167
TM
88
23
HM1.24
P3.19
IS-1
IS-2
IS-3
IS-4
IS-5

Fig.16

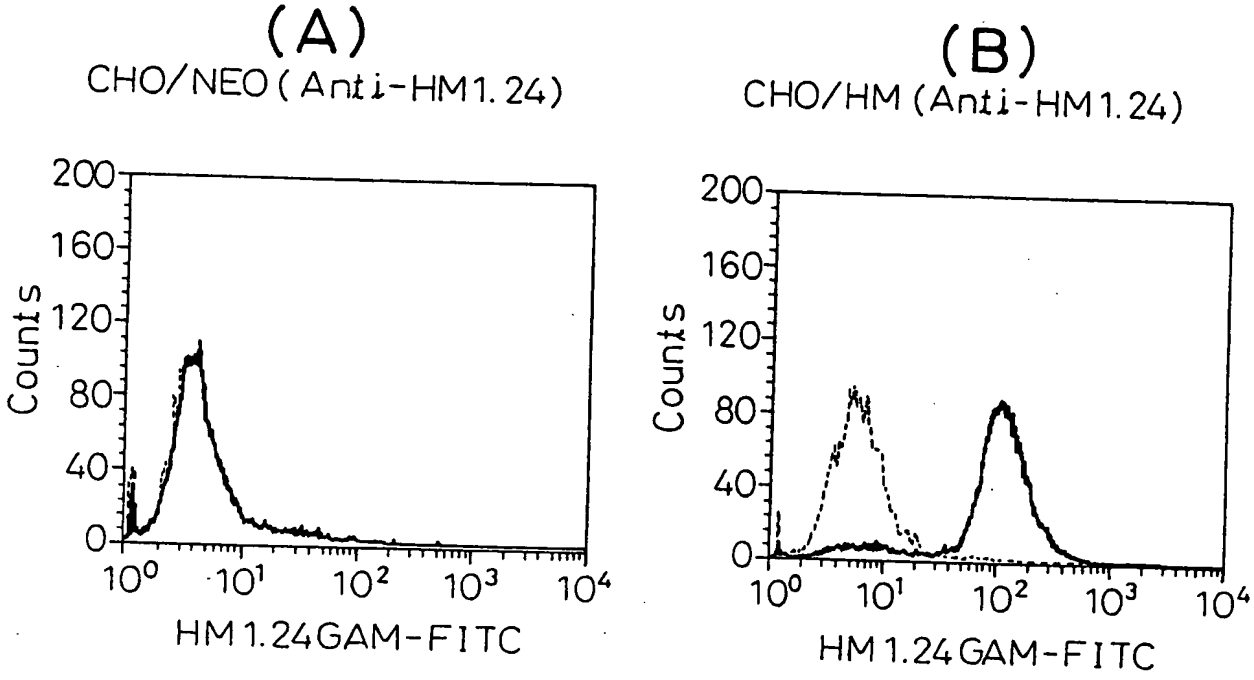
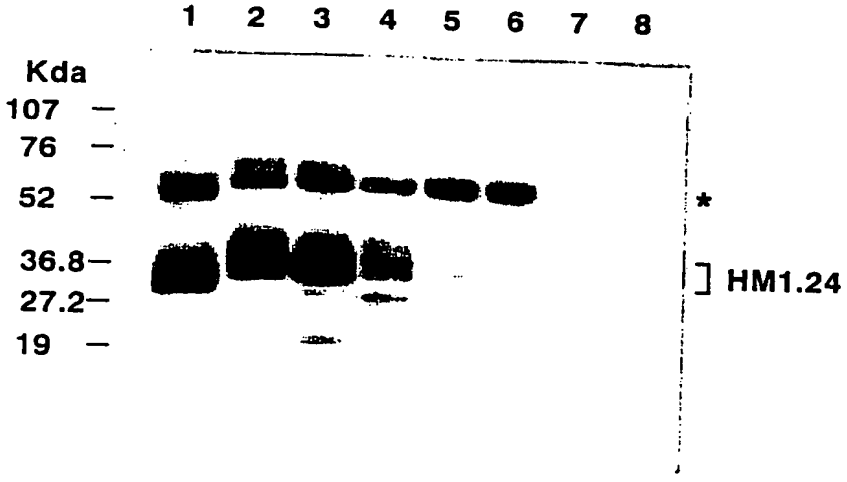
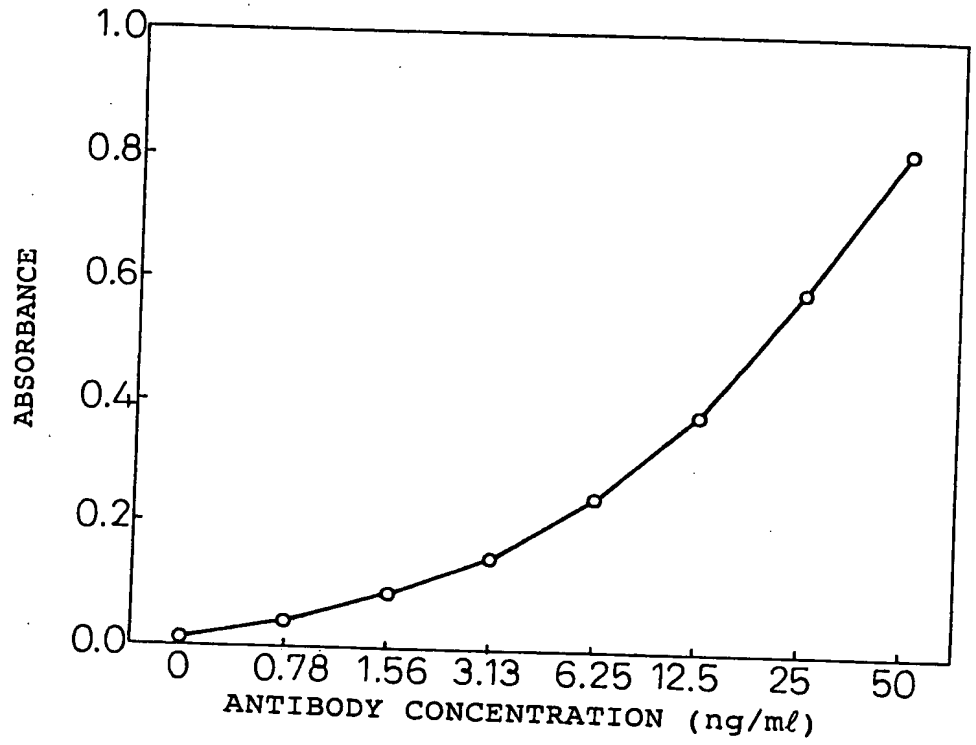


Fig.17



LANE 1: KPMM2 (EQUIVALENT TO 5 × 10^5 CELLS)
LANE 2: RPMI8226 (25 × 10^5 CELLS)
LANE 3: U266 (25 × 10^5 CELLS)
LANE 4: CHO/HM (5 × 10^5 CELLS)
LANE 5: CHO/NEO (5 × 10^5 CELLS)
LANE 6: NONE
LANE 7: KPMM2 (5 x 10^5 CELLS)

Fig.18


1.0
0.8
0.6
0.4
0.2
0.0
ABSORBANCE
0
0.78
1.56
3.13
6.25
12.5
25
50
ANTIBODY CONCENTRATION (ng/mℓ)